

IN THE SPECIFICATION

Please amend paragraph [0027] starting at the bottom of page 13 as follows:

[0027]

The names, numbers, and functions of respective genes in the Tables contained herein conform with the *Bacillus subtilis* genome data reported in Nature, 390, 249 to 256 (1997) and made public by JAFAN (Japan Functional Analysis Network for *Bacillus subtilis*; BSORF DB) on the Internet by inserting (~~http://~~*bacillus.genome.ad.jp/* renewed June 17, 2003) after the hypertext transfer protocol *http://*.

Please amend paragraph [0045] on page 23 of the specification as follows:

[0045]

The promoter sequence employed in the present invention which is recognized by a sigma factor relating specifically to transcription during the sporulation stage includes promoter sequences for expressing ortholog genes of a gene of *Bacillus subtilis* or each of some genes constituting an operon of *Bacillus subtilis*, listed in Tables 2 and 3. The ortholog genes are preferably derived from bacteria belonging to the genus *Bacillus*. The ortholog genes can be located by use of a Create/view Orthologous gene table program of Microbial Genome Database (MBGD, ~~http://~~ available by inserting *mbgd.genome.ad.jp/* after the hypertext transfer protocol *http://*) published on the internet. Examples of the ortholog genes of the *sigH* gene of *Bacillus subtilis* include a *sigH* (BH0115) gene of *Bacillus halodulans* and a BC0114 gene of *Bacillus cereus*. Examples of the ortholog genes of each of some genes constituting the *spoIIA* operon of *Bacillus subtilis* include a *sigF* (BH1538) gene, a *spoIIAB* (BH1537) gene, and a *spoIIAA* (BH1536) gene, these three genes being of *Bacillus halodulans*; and a BC4072 gene, a BC4073 gene, and a BC4074 gene, these three genes being of *Bacillus cereus*.